

SEQUENCE LISTING

<110> Cannon, Paul David
Sankuratri, Suryanarayana

<120> Human Intestinal Npt2B

<130> ROCH-001

<150> 60/119,321

<151> 1999-02-09

<160> 2

<170> FastSEQ for Windows Version 4.0

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FIG. 1

1 MAPWPELGDA QPNPDKYLEG AAGQQPTAPD KSKETNKNNT EAPVTKIELL
51 PSYSTATLID EPTEVDDPWN LPTLQDSGIK WSERDTKGKI LCFFQGIGRL
101 ILLLGFLYFF VCSLDILSSA FQLVGGKMAG QFFSNSSIMS NPLLGLVIGV
151 LVTVLVQSSS TSTSIVVSMV SSSLLTVRAA IPIIMGANIG TSITNTIVAL
201 MQVGDRSEFR RAFAGATVHD FFNWLSVLVL LPVEVATHYL EIITQLIVES
251 FHFKNGEDAP DLLKVITKPF TKLIVQLDKK VISQIAMNDE KAKNKSLVKI
301 WCKTFTNKTQ INVTVPSTAN CTSPSLCWTD GIQNWTKMNV TYKENIAKCQ
351 HIFVNFHLPD LAVGTILLIL SLLVLCGCLI MIVKILGSVL KGQVATVIKK
401 TINTDFPPFF AWLTGYLAIL VGAGMTFIVQ SSSVFTSALT PLIGIGVITI
451 ERAYPLTLGS NIGTTTTAIL AALASPGNAL RSSLQIALCH FFFNISGILL
501 WYPIPFTRLP IRMAKGLGNI SAKYRWFAVF YLIIFFFLIP LTVFGLSLAG
551 WRVLVGVGVP VVFIIILVLC LRLQSRCPR VLPKKLQNNW FLPLWMRSK
601 PWDVVSKFT GCFQMRCCCC CRVCCRACCL LCGCPKCCRC SKCCEDLEEA
651 QEGQDVPVKA PETFDNITIS REAQGEVPAS DSKTECTAL* (SEQ ID NO:01)

FIG. 2i

1 CTGACGTAGG CCCAGCACCT GCGGAGGGAG CGCTGACCAT GGCTCCCTGG
51 CCTGAATTGG GAGATGCCCA GCCCAACCCC GATAAGTACC TCGAAGGGGC
101 CGCAGGTCAG CAGCCCACTG CCCCTGATAA AAGCAAAGAG ACCAACAAAA
151 ATAACACTGA GGCACCTGTA ACCAAGATTG AACTTCTGCC GTCCTACTCC
201 ACGGCTACAC TGATAGATGA GCCCACTGAG GTGGATGACC CCTGGAACCT
251 ACCCACTCTT CAGGACTCGG GGATCAAGTG GTCAGAGAGA GACACCAAAG
301 GGAAGATTCT CTGTTTCTTC CAAGGGATTG GGAGATTGAT TTTACTTCTC
351 GGATTTCTCT ACTTTTTCGT GTGCTCCCTG GATATTCTTA GTAGCGCCTT
401 CCAGCTGGTT GGAGGAAAAA TGGCAGGACA GTTCTTCAGC AACAGCTCTA
451 TTATGTCCAA CCCTTTGTTG GGGCTGGTGA TCGGGGTGCT GGTGACCGTC
501 TTGGTGCAGA GCTCCAGCAC CTCAACGTCC ATCGTTGTCA GCATGGTGTC
551 CTCTTCATTG CTCACTGTTC GGGCTGCCAT CCCCATTATC ATGGGGGGCCA
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651 GATCGGAGTG AGTTCAGAAG AGCTTTTGCA GGAGCCACTG TCCATGACTT
701 CTTCAACTGG CTGTCCGTGT TGGTGCTCTT GCCCGTGGAG GTGGCCACCC
751 ATTACCTCGA GATCATAACC CAGCTTATAG TGGAGAGCTT CCACTTCAAG
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851 AAAGCTCATT GTCCAGCTGG ATAAAAAAGT TATCAGCCAA ATTGCAATGA
901 ACGATGAAAA AGCGAAAAAC AAGAGTCTTG TCAAGATTG GTGCAAAACT
951 TTTACCAACA AGACCCAGAT TAACGTCACT GTTCCCTCGA CTGCTAACTG
1001 CACCTCCCTT TCCCTCTGTT GGACGGATGG CATCCAAAAC TGGACCATGA
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1101 AATTTCCACC TCCCGGATCT TGCTGTGGGC ACCATCTTGC TCATACTCTC
1151 CCTGCTGGTC CTCTGTGGTT GCCTGATCAT GATTGTCAAG ATCCTGGGCT
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1701 GTTGGTGTG GGGTTCCCGT CGTCTTCATC ATCATCCTGG TACTGTGCCT
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2201 GATATTGCT CCCCATTAGC GAATGAAATT GATGCAGTCC TACCTAACTC
2251 GATTCCCTTT GGCTTGGTGG GTAGGCCTGC AGGGCACTTT TATTCCAACC
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2451 AAAGAAAAGG CCCAGGGAAG GAATGTATGA GAGGCTCTCC CAGATGAGGA
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2601 AGCCTGGGTC AGGGGACATA GTGTCATTGT TTGGAAACTG CAGACCACAA

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FIG. 2ii

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2751 TTGTCCTCAT GCTTCGGGGA TGGGAGCCAC GCCTGAACTA GAGTTCAGGC
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